

Sequence Listing.ST25.txt  
SEQUENCE LISTING

&lt;110&gt; Novartis AG

&lt;120&gt; Use of fibroblast growth factor fragments

&lt;130&gt; 4-33264A

&lt;160&gt; 4

&lt;170&gt; PatentIn version 3.1

&lt;210&gt; 1

&lt;211&gt; 251

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 1.

Met Leu Gly Ala Arg Leu Arg Leu Trp Val Cys Ala Leu Cys Ser Val  
1 5 10 15Cys Ser Met Ser Val Leu Arg Ala Tyr Pro Asn Ala Ser Pro Leu Leu  
20 25 30Gly Ser Ser Trp Gly Gly Leu Ile His Leu Tyr Thr Ala Thr Ala Arg  
35 40 45Asn Ser Tyr His Leu Gln Ile His Lys Asn Gly His Val Asp Gly Ala  
50 55 60Pro His Gln Thr Ile Tyr Ser Ala Leu Met Ile Arg Ser Glu Asp Ala  
65 70 75 80Gly Phe Val Val Ile Thr Gly Val Met Ser Arg Arg Tyr Leu Cys Met  
85 90 95Asp Phe Arg Gly Asn Ile Phe Gly Ser His Tyr Phe Asp Pro Glu Asn  
100 105 110Cys Arg Phe Gln His Gln Thr Leu Glu Asn Gly Tyr Asp Val Tyr His  
115 120 125Ser Pro Gln Tyr His Phe Leu Val Ser Leu Gly Arg Ala Lys Arg Ala  
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130 135 140

Phe Leu Pro Gly Met Asn Pro Pro Pro Tyr Ser Gln Phe Leu Ser Arg  
145 150 155 160

Arg Asn Glu Ile Pro Leu Ile His Phe Asn Thr Pro Ile Pro Arg Arg  
165 170 175

His Thr Arg Ser Ala Glu Asp Asp Ser Glu Arg Asp Pro Leu Asn Val  
180 185 190

Leu Lys Pro Arg Ala Arg Met Thr Pro Ala Pro Ala Ser Cys Ser Gln  
195 200 205

Glu Leu Pro Ser Ala Glu Asp Asn Ser Pro Met Ala Ser Asp Pro Leu  
210 215 220

Gly Val Val Arg Gly Gly Arg Val Asn Thr His Ala Gly Gly Thr Gly  
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Pro Glu Gly Cys Arg Pro Phe Ala Lys Phe Ile  
245 250

<210> 2

<211> 75

<212> PRT

<213> Homo sapiens

<400> 2

His Thr Arg Ser Ala Glu Asp Asp Ser Glu Arg Asp Pro Leu Asn Val  
1 5 10 15

Leu Lys Pro Arg Ala Arg Met Thr Pro Ala Pro Ala Ser Cys Ser Gln  
20 25 30

Glu Leu Pro Ser Ala Glu Asp Asn Ser Pro Met Ala Ser Asp Pro Leu  
35 40 45

Gly Val Val Arg Gly Gly Arg Val Asn Thr His Ala Gly Gly Thr Gly  
50 55 60

Pro Glu Gly Cys Arg Pro Phe Ala Lys Phe Ile  
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<210> 3

<211> 756

<212> DNA

## Sequence Listing.ST25.txt

&lt;213&gt; Homo sapiens

&lt;400&gt; 3

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gtggatggcg caccatca gaccatctac agtgccctga tgatcagatc agaggatgct    240
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gccgaggacg actcggagcg ggacccccctg aacgtgctga agccccgggc ccgatgacc    600
ccggccccgg cctcctgttc acaggagctc ccgagcgccg aggacaacag cccgatggcc    660
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&lt;210&gt; 4

&lt;211&gt; 228

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 4

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agccccgatgg ccagtgaccc attaggggtg gtcaggggcg gtcgagtga cgcgcacgct    180
gggggaacgg gcccggaagg ctgccgcccc ttcgccaagt tcattctag    228
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